Attorney Docket No. Application No. 10/579.780 042715-5019 ORMATION DISCLOSURE CITATION e several sheets if necessary) Applicants: Yoshiva ODA et al. Page 1 of 2 JUN 2 3- 2006 PTO Form 1449 Filing Date: May 18, 2006 Group Art Unit: Unassigned June 23, 2006 U.S. PATENT DOCUMENTS Class Sub-Filing Date Initial Document No. Date Name Class US 6.391.649 B1 May 21, 2002 Chait et al 436 173 May 4, 1999 1. FOREIGN PATENT DOCUMENTS Translation Document No. Date Country Class Sub-Class 2 WO 03/016861 02/27/2003 PCT Abstract 3 IP P 2003-107066 A 09/04/2003 Japan IP P 2000-131305 A 12/05/2000 Japan Abstract OTHER DOCUMENTS (Including Author, Title, Date, Pertinent Pages, etc.) "Post-Genomic Mass Spectrometry" 1993, pp. 39-72, Toshimitsu Niwa Ed., Kagakudojin (English translation of abstract enclosed) Carney et al. "De novo peptide sequencing and quantitative profiling of complex protein mixtures using masscoded abundance tagging" Nat Biotechnol, 2002 Feb; 20(2):163-170 7 Drewes et al. "Global approaches to protein-protein interactions" Curr Opin Cell Biol. 2003 Apr;15(2):199-205 Gavin et al. "Protein complexes and proteome organization from yeast to man" Curr Opin Chem Biol. 2003 Feb;7(1):21-27 Gerber et al. "Absolute quantification of proteins and phosphoproteins from cell lysates by tandem MS" Proc Natl Acad Sci U S A. 2003 Jun 10;100(12):6940-6945. Epub 2003 May 27 Goodlett et al. "Differential stable isotope labeling of peptides for quantitation and de novo sequence derivation" Rapid Commun Mass Spectrom. 2001;15(14):1214-1221 Griffin et al. "Toward a high-throughput approach to quantitative proteomic analysis: expression-dependent protein 11. identification by mass spectrometry" J Am Soc Mass Spectrom. 2001 Dec;12(12):1238-1246 Gygi et al. "Quantitative analysis of complex protein mixtures using isotope-coded affinity tags" Nat Biotechnol. 12. 1999 Oct;17(10):994-999 Han et al. "Quantitative profiling of differentiation-induced microsomal proteins using isotope-coded affinity tags 13. and mass spectrometry" Nat Biotechnol, 2001 Oct;19(10):946-951 Ishihama et al. "Simple and sensitive quantitation method for mevalonic acid in plasma using gas 14 chromatography/mass spectrometry" Rapid Commun Mass Spectrom. 1994 May;8(5):377-380 Kahn "From genome to proteome: looking at a cell's proteins" Science. 1995 Oct 20;270(5235):369-370 15. Matsui et al. "Direct determination of E2020 enantiomers in plasma by liquid chromatography-mass spectrometry and column-switching techniques" J Chromatogr A. 1995 Mar 3;694(1):209-218. Matsui et al. "Simultaneous determination of done pezil (aricept) enantiomers in human plasma by liquid 17. chromatography-electrospray tandem mass spectrometry" J Chromatogr B Biomed Sci Appl. 1999 Jun 11;729(1-2):147-155 Neubauer et al. "Mass spectrometry and EST-database searching allows characterization of the multi-protein spliceosome complex" Nat Genet. 1998 Sep;20(1):46-50 Oda et al. "Accurate quantitation of protein expression and site-specific phosphorylation" Proc Natl Acad Sci U S A. 1999 Jun 8:96(12):6591-6596 Examiner Date Considered 08/24/2009 /Yelena Gakh/ Examiner: Initial if reference considered, whether or not citation is in conformance with MPEP 609; draw line through citation if not

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